

=====

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=10; day=29; hr=14; min=12; sec=15; ms=896;  
]

=====

\*\*\*\*\*

Reviewer Comments:

<110> APPLICANT: Hakoto KOBAYAHSI

Yugo HABATA

Ryo FUJII

Shuji HINUMA

<120> TITLE OF INVENTION: Methods of Screening for Ligands for FPRL2

<130> FILE REFERENCE: 3171 US0P

<140> CURRENT APPLICATION NUMBER:10554234

<141> CURRENT FILING DATE:2005-10-21

<150> PRIOR APPLICATION NUMBER: PCT/JP2004/005829

<151> PRIOR FILING DATE: 2004-04-22

<150> PRIOR APPLICATION NUMBER: JP 2003-118760

<151> PRIOR FILING DATE: 2003-04-23

<160> NUMBER OF SEQ ID NOS: 8

<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

Please do not insert alpha numeric headings. Please make necessary changes. This type of error is seen globally throughout the sequence listing.

\*\*\*\*\*

Application No: 10554234 Version No: 2.0

**Input Set:**

**Output Set:**

**Started:** 2008-09-30 12:20:13.960  
**Finished:** 2008-09-30 12:20:14.257  
**Elapsed:** 0 hr(s) 0 min(s) 0 sec(s) 297 ms  
**Total Warnings:** 3  
**Total Errors:** 0  
**No. of SeqIDs Defined:** 8  
**Actual SeqID Count:** 8

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 333	tabs used in amino acid numbering SEQID (8)

<110> APPLICANT: Hakoto KOBAYAHSI

Yugo HABATA

Ryo FUJII

Shuji HINUMA

<120> TITLE OF INVENTION: Methods of Screening for Ligands for FPRL2

<130> FILE REFERENCE: 3171 US0P

<140> CURRENT APPLICATION NUMBER: 10554234

<141> CURRENT FILING DATE: 2005-10-21

<150> PRIOR APPLICATION NUMBER: PCT/JP2004/005829

<151> PRIOR FILING DATE: 2004-04-22

<150> PRIOR APPLICATION NUMBER: JP 2003-118760

<151> PRIOR FILING DATE: 2003-04-23

<160> NUMBER OF SEQ ID NOS: 8

<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

Met Glu Thr Asn Phe Ser Ile Pro Leu Asn Glu Thr Glu Glu Val Leu  
5 10 15  
Pro Glu Pro Ala Gly His Thr Val Leu Trp Ile Phe Ser Leu Leu Val  
20 25 30  
His Gly Val Thr Phe Val Phe Gly Val Leu Gly Asn Gly Leu Val Ile  
35 40 45  
Trp Val Ala Gly Phe Arg Met Thr Arg Thr Val Asn Thr Ile Cys Tyr  
50 55 60  
Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Ser Ala Ile Leu Pro Phe  
65 70 75 80  
Arg Met Val Ser Val Ala Met Arg Glu Lys Trp Pro Phe Ala Ser Phe  
85 90 95  
Leu Cys Lys Leu Val His Val Met Ile Asp Ile Asn Leu Phe Val Ser  
100 105 110  
Val Tyr Leu Ile Thr Ile Ile Ala Leu Asp Arg Cys Ile Cys Val Leu  
115 120 125  
His Pro Ala Trp Ala Gln Asn His Arg Thr Met Ser Leu Ala Lys Arg  
130 135 140  
Val Met Thr Gly Leu Trp Ile Phe Thr Ile Val Leu Thr Leu Pro Asn  
145 150 155 160  
Phe Ile Phe Trp Thr Thr Ile Ser Thr Thr Asn Gly Asp Thr Tyr Cys  
165 170 175  
Ile Phe Asn Phe Ala Phe Trp Gly Asp Thr Ala Val Glu Arg Leu Asn  
180 185 190  
Val Phe Ile Thr Met Ala Lys Val Phe Leu Ile Leu His Phe Ile Ile  
195 200 205  
Gly Phe Thr Val Pro Met Ser Ile Ile Thr Val Cys Tyr Gly Ile Ile  
210 215 220  
Ala Ala Lys Ile His Arg Asn His Met Ile Lys Ser Ser Arg Pro Leu  
225 230 235 240  
Arg Val Phe Ala Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro  
245 250 255  
Tyr Glu Leu Ile Gly Ile Leu Met Ala Val Trp Leu Lys Glu Met Leu  
260 265 270  
Leu Asn Gly Lys Tyr Lys Ile Ile Leu Val Leu Ile Asn Pro Thr Ser  
275 280 285  
Ser Leu Ala Phe Phe Asn Ser Cys Leu Asn Pro Ile Leu Tyr Val Phe

290	295	300
Met	Gly Arg Asn Phe Gln Glu Arg Leu Ile Arg Ser Leu Pro Thr Ser	
305	310	315
Leu Glu Arg Ala Leu Thr Glu Val Pro Asp Ser Ala Gln Thr Ser Asn		
325	330	335
Thr His Thr Thr Ser Ala Ser Pro Pro Glu Glu Thr Glu Leu Gln Ala		
340	345	350
Met		

<210> SEQ ID NO 2

<211> LENGTH: 1059

<212> TYPE: DNA

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 2

atggaaacca	acttctccat	tcctctgaat	gaaaactgagg	aggtgctccc	tgagcctgct	60
ggccacaccg	ttctgtggat	cttctcatttgc	ctagtccacg	gagtcaccc	tgtcttcggg	120
gtcctgggca	atgggcttgt	gatctgggtg	gctggattcc	ggatgacacg	cacagtcaac	180
accatctgtt	acctgaacct	ggcccttagct	gacttctctt	tcagtgccat	cctaccattc	240
cgaatggtct	cagtcgccccat	gagagaaaaaa	tggccttttgc	cgtcattccct	atgttaagtta	300
gttcatgtt	tgtatagacat	caacctgttt	gtcagtgct	acctgatcac	catcatgtct	360
ctggaccgct	gtatattgtgt	cctgcattcca	gcctggggcc	agaaccatcg	caccatgagt	420
ctggccaaga	gggtgatgac	gggactctgg	atttcacca	tagtccttac	cttaccaaata	480
ttcatcttct	ggactacaat	aagtactacg	aatggggaca	catactgtat	tttcaacttt	540
gcattctggg	gtgacactgc	tgttagagagg	ttgaacgtgt	tcattaccat	ggccaagggtc	600
tttctgatcc	tccacttcat	tattggcttc	acgggtgccta	tgtccatcat	cacagtctgc	660
tatggatca	tcgctgccaa	aattcacaga	aaccacatga	ttaaatccag	ccgtccctta	720
cgtgtcttcg	ctgctgtggt	ggcttcttc	ttcatctgtt	ggttccctta	tgaactaatt	780
ggcattctaa	tggcagtctg	gctcaaagag	atgttggtaa	atggcaaata	caaaatcatt	840
cttgcctga	ttaacccaaac	aagctccttg	gccttttta	acagctgcct	caacccaatt	900
ctctacgtct	ttatgggtcg	taacttccaa	gaaagactga	ttcgctctt	gcccactagt	960
ttggagaggg	ccctgactga	ggtccctgac	tcagcccaga	ccagcaacac	acacaccact	1020
tctgcttac	ctcctgagga	gacggagtt	caagcaatg			1059

<210> SEQ ID NO 3

<211> LENGTH: 6

<212> TYPE: PRT

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: amino acid sequence of GHRP-6

<220> FEATURE:

<223> OTHER INFORMATION: Trp is a D-form

<400> SEQUENCE: 3

His	Trp	Ala	Trp	Phe	Lys
-----	-----	-----	-----	-----	-----

1		5
---	--	---

<210> SEQ ID NO 4

<211> LENGTH: 11

<212> TYPE: PRT

<213> ORGANISM: Aplysia sp.

<400> SEQUENCE: 4

Ala	Arg	Pro	Gly	Tyr	Leu	Ala	Phe	Pro	Arg	Met
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

1		5		10
---	--	---	--	----

<210> SEQ ID NO 5

<211> LENGTH: 12

<212> TYPE: PRT  
<213> ORGANISM: Sus scrofa  
<400> SEQUENCE: 5  
Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe  
1 5 10

<210> SEQ ID NO 6  
<211> LENGTH: 36  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 6  
Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp  
1 5 10 15  
Leu Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr  
20 25 30  
Arg Gln Arg Tyr  
35

<210> SEQ ID NO 7  
<211> LENGTH: 10  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 7  
Gly Asn His Trp Ala Val Gly His Leu Met  
1 5 10

<210> SEQ ID NO 8  
<211> LENGTH: 6  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 8  
Met Val Met Tyr Lys Trp  
1 5